CRFI #9



OIPE

RAW SEQUENCE LISTING DATE: 07/17/2002 PATENT APPLICATION: US/10/001,934 TIME: 07:14:31

Input Set : N:\Crf3\Datahold\EFS\10001934\GPCG-P01-003SequenceListing.APP

Output Set: N:\CRF3\07172002\J001934.raw

3 <110> APPLICANT: NAGY, ZOLTAN

```
5 <120> TITLE OF INVENTION: HUMAN PEPTIDES/PROTEINS CAUSING OR LEADING TO THE
             KILLING OF CELLS INCLUDING LYMPHOID TUMOR CELLS
     8 <130> FILE REFERENCE: GPCG-P01-003
     10 <140> CURRENT APPLICATION NUMBER: 10/001,934
    11 <141> CURRENT FILING DATE: 2001-11-15
    13 <160> NUMBER OF SEQ ID NOS: 63
    15 <170> SOFTWARE: PatentIn Ver. 2.1
    17 <210> SEQ ID NO: 1
    18 <211> LENGTH: 10
    19 <212> TYPE: PRT
    20 <213> ORGANISM: Artificial Sequence
     22 <220> FEATURE:
     23 <223> OTHER INFORMATION: Description of Artificial Sequence: CDR3 consensus
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     26 <220> FEATURE:
     27 <221> NAME/KEY: MOD_RES
     28 <222> LOCATION: (1)..(4)
     29 <223> OTHER INFORMATION: Any amino acid
     31 <220> FEATURE:
     32 <221> NAME/KEY: MOD_RES
     33 <222> LOCATION: (7)
     34 <223> OTHER INFORMATION: Any amino acid
     36 <220> FEATURE:
     37 <221> NAME/KEY: MOD_RES
     38 <222> LOCATION: (10)
     39 <223> OTHER INFORMATION: Any amino acid
     41 <400> SEQUENCE: 1
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     43
     46 <210> SEQ ID NO: 2
     47 <211> LENGTH: 8
     48 <212> TYPE: PRT
     49 <213> ORGANISM: Artificial Sequence
     51 <220> FEATURE:
     52 <223> OTHER INFORMATION: Description of Artificial Sequence: CDR3 consensus
              sequence
     55 <220> FEATURE:
     56 <221> NAME/KEY: MOD_RES
     57 <222> LOCATION: (5)..(8)
     58 <223> OTHER INFORMATION: Any amino acid
     60 <400> SEQUENCE: 2
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W--> 61 Gln Ser Tyr Asp Xaa Xaa Xaa Xaa

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     65 <210> SEQ ID NO: 3
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     67 <212> TYPE: PRT
     68 <213> ORGANISM: Homo sapiens
     70 <400> SEQUENCE: 3
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     72
                          5
          1
     75 <210> SEQ ID NO: 4
     76 <211> LENGTH: 8
     77 <212> TYPE: PRT
     78 <213> ORGANISM: Homo sapiens
     80 <400> SEQUENCE: 4
     81 Gln Ser Tyr Asp Leu Ile Arg His
     85 <210> SEQ ID NO: 5
     86 <211> LENGTH: 8
     87 <212> TYPE: PRT
     88 <213> ORGANISM: Homo sapiens
     90 <400> SEQUENCE: 5
     91 Gln Ser Tyr Asp Met Asn Val His
          1
     95 <210> SEQ ID NO: 6
     96 <211> LENGTH: 13
     97 <212> TYPE: PRT
     98 <213> ORGANISM: Artificial Sequence
     100 <220> FEATURE:
     101 <223> OTHER INFORMATION: Description of Artificial Sequence: CDR1 consensus
     102
               sequence
     104 <220> FEATURE:
     105 <221> NAME/KEY: MOD_RES
     106 <222> LOCATION: (4)..(5)
     107 <223> OTHER INFORMATION: Any amino acid
     109 <220> FEATURE:
    110 <221> NAME/KEY: MOD_RES
     111 <222> LOCATION: (9)
     112 <223> OTHER INFORMATION: Any amino acid
     114 <220> FEATURE:
     115 <221> NAME/KEY: MOD_RES
     116 <222> LOCATION: (13)
     117 <223> OTHER INFORMATION: Any amino acid
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     124 <210> SEQ ID NO: 7
     125 <211> LENGTH: 13
     126 <212> TYPE: PRT
     127 <213> ORGANISM: Artificial Sequence
     129 <220> FEATURE:
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130 <223> OTHER INFORMATION: Description of Artificial Sequence: CDR1 consensus
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138 <210> SEQ ID NO: 8
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140 <212> TYPE: PRT
141 <213> ORGANISM: Artificial Sequence
143 <220> FEATURE:
144 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
145
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147 <400> SEQUENCE: 8
148 Trp Ser His Pro Gln Phe Glu Lys
149
152 <210> SEQ ID NO: 9
153 <211> LENGTH: 4
154 <212> TYPE: PRT
155 <213> ORGANISM: Artificial Sequence
157 <220> FEATURE:
158 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic FLAG
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162 Asp Tyr Lys Asp
163
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167 <211> LENGTH: 17
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169 <213> ORGANISM: Artificial Sequence
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172 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
174 <400> SEQUENCE: 10
175 gtggtggttc cgatatc
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178 <210> SEQ ID NO: 11
179 <211> LENGTH: 44
180 <212> TYPE: DNA
181 <213> ORGANISM: Artificial Sequence
183 <220> FEATURE:
184 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
186 <400> SEQUENCE: 11
187 agcgtcacac tcggtgcggc tttcggctgg ccaagaacgg gtta
                                                                       44
190 <210> SEQ ID NO: 12
191 <211> LENGTH: 13
192 <212> TYPE: PRT
193 <213> ORGANISM: Homo sapiens
195 <400> SEQUENCE: 12
196 Ser Gly Ser Ser Ser Asn Ile Gly Ser Asn Tyr Val Ser
197
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200 <210> SEQ ID NO: 13
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211 <211> LENGTH: 13
212 <212> TYPE: PRT
213 <213> ORGANISM: Homo sapiens
215 <400> SEQUENCE: 14
216 Ser Gly Ser Glu Ser Asn Ile Gly Ser Asn Tyr Val Ala
217 1
                      5
220 <210> SEQ ID NO: 15
221 <211> LENGTH: 13
222 <212> TYPE: PRT
223 <213> ORGANISM: Homo sapiens
225 <400> SEQUENCE: 15
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230 <210> SEQ ID NO: 16
231 <211> LENGTH: 13
232 <212> TYPE: PRT
233 <213> ORGANISM: Homo sapiens
235 <400> SEQUENCE: 16
236 Ser Gly Ser Glu Pro Asn Ile Gly Ser Asn Tyr Val Phe
237 1
240 <210> SEQ ID NO: 17
241 <211> LENGTH: 13
242 <212> TYPE: PRT
243 <213> ORGANISM: Homo sapiens
245 <400> SEQUENCE: 17
246 Ser Gly Ser Glu Ser Asn Ile Gly Asn Asn Tyr Val Gly
247
250 <210> SEQ ID NO: 18
251 <211> LENGTH: 13
252 <212> TYPE: PRT
253 <213> ORGANISM: Homo sapiens
255 <400> SEQUENCE: 18
256 Ser Gly Ser Glu Ser Asn Ile Gly Ala Asn Tyr Val Asn
257
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260 <210> SEQ ID NO: 19
261 <211> LENGTH: 10
262 <212> TYPE: PRT
263 <213> ORGANISM: Homo sapiens
265 <400> SEQUENCE: 19
266 Gln Tyr Gly His Arg Gly Gly Phe Asp His
270 <210> SEQ ID NO: 20
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Output Set: N:\CRF3\07172002\J001934.raw

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271 <211> LENGTH: 9
272 <212> TYPE: PRT
273 <213> ORGANISM: Homo sapiens
275 <400> SEQUENCE: 20
276 Gly Tyr Gly Arg Tyr Ser Pro Asp Leu
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280 <210> SEQ ID NO: 21
281 <211> LENGTH: 8
282 <212> TYPE: PRT
283 <213> ORGANISM: Homo sapiens
285 <400> SEQUENCE: 21
286 Gln Gln Tyr Ser Asn Leu Pro Phe
287
290 <210> SEQ ID NO: 22
291 <211> LENGTH: 8
292 <212> TYPE: PRT
293 <213> ORGANISM: Homo sapiens
295 <400> SEQUENCE: 22
296 Gln Ser Tyr Asp Met Pro Gln Ala
297
300 <210> SEQ ID NO: 23
301 <211> LENGTH: 8
302 <212> TYPE: PRT
303 <213> ORGANISM: Homo sapiens
305 <400> SEQUENCE: 23
306 Gln Ser Tyr Asp Leu Thr Met Gly
307
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310 <210> SEQ ID NO: 24
311 <211> LENGTH: 8
312 <212> TYPE: PRT
313 <213> ORGANISM: Homo sapiens
315 <400> SEQUENCE: 24
316 Gln Ser Tyr Asp Phe Ser His Tyr
317
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320 <210> SEQ ID NO: 25
321 <211> LENGTH: 8
322 <212> TYPE: PRT
323 <213> ORGANISM: Homo sapiens
325 <400> SEQUENCE: 25
326 Gln Ser Tyr Asp Ile Gln Leu His
327
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330 <210> SEQ ID NO: 26
331 <211> LENGTH: 8
332 <212> TYPE: PRT
333 <213> ORGANISM: Homo sapiens
335 <400> SEQUENCE: 26
336 Gln Ser Tyr Asp Phe Ser Val Tyr
      1
340 <210> SEQ ID NO: 27
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RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/001,934

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Input Set : N:\Crf3\Datahold\EFS\10001934\GPCG-P01-003SequenceListing.APP

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 1,2,3,4,7,10

Seq#:2; Xaa Pos. 5,6,7,8

Seq#:6; Xaa Pos. 4,5,9,13

VERIFICATION SUMMARY
PATENT APPLICATION: US/10/001,934

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Output Set: N:\CRF3\07172002\J001934.raw

L:42 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0 L:61 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0 L:120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0